### **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/583.084
Source:	IFWP.
Date Processed by STIC:	6/28/06
· ·	

## ENTERED

## CRF Errors Edited by the STIC Systems Branch

Serial	Number: 10/583, 084	CRF Edit Date: _ Edited by:	6/28/ 
	Realigned nucleic acid/amino acid numbers/text text "wrapped" to the next line	in cases where the	e sequence
	Corrected the SEQ ID NO. Sequence numbers e	edited were:	
	Inserted or corrected a nucleic number at the en NO's edited:	d of a nucleic line.	SEQ ID
_	Deleted: invalid beginning/end-of-file text;	page numbers	
	Inserted mandatory headings/numeric identifier	s, specifically:	
	Moved responses to same line as heading/numer	ic identifier, specif	ically:
	Other:	·	

Revised 09/09/2003



IFWP

RAW SEQUENCE LISTING DATE: 06/28/2006
PATENT APPLICATION: US/10/583,084 TIME: 10:45:12

Input Set : A:\PTO.AMC.txt

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3 <110> APPLICANT: SUNTORY LIMITED
     5 <120> TITLE OF INVENTION: Arachidonic acid-containing plants and use of the plants
     7 <130> FILE REFERENCE: SU0423
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/583,084
C--> 10 <141> CURRENT FILING DATE: 2006-06-15
     12 <150> PRIOR APPLICATION NUMBER: JP 2003-419124
     13 <151> PRIOR FILING DATE: 2003-12-17
     15 <150> PRIOR APPLICATION NUMBER: JP 2004-097089
     16 <151> PRIOR FILING DATE: 2004-03-29
     18 <160> NUMBER OF SEQ ID NOS: 28
     20 <170> SOFTWARE: PatentIn Ver. 2.1
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 457
     24 <212> TYPE: PRT
     25 <213> ORGANISM: Mortierella alpina
     27 <400> SEQUENCE: 1
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                                             10
     31 Asn Ala Glu Ala Leu Asn Glu Gly Lys Lys Asp Ala Glu Ala Pro Phe
     34 Leu Met Ile Ile Asp Asn Lys Val Tyr Asp Val Arg Glu Phe Val Pro
     37 Asp His Pro Gly Gly Ser Val Ile Leu Thr His Val Gly Lys Asp Gly
                                 55
     40 Thr Asp Val Phe Asp Thr Phe His Pro Glu Ala Ala Trp Glu Thr Leu
                             70
     43 Ala Asn Phe Tyr Val Gly Asp Ile Asp Glu Ser Asp Arg Ala Ile Lys
                        85
                                             90
     46 Asn Asp Asp Phe Ala Ala Glu Val Arg Lys Leu Arg Thr Leu Phe Gln
                                        105
     49 Ser Leu Gly Tyr Tyr Asp Ser Ser Lys Ala Tyr Tyr Ala Phe Lys Val
     50
                                    120
    52 Ser Phe Asn Leu Cys Ile Trp Gly Leu Ser Thr Phe Ile Val Ala Lys
                                135
                                                    140
     55 Trp Gly Gln Thr Ser Thr Leu Ala Asn Val Leu Ser Ala Ala Leu Leu
                            150
     58 Gly Leu Phe Trp Gln Gln Cys Gly Trp Leu Ala His Asp Phe Leu His
                                            170
                        165
     61 His Gln Val Phe Gln Asp Arg Phe Trp Gly Asp Leu Phe Gly Ala Phe
                   180
                                        185
    64 Leu Gly Gly Val Cys Gln Gly Phe Ser Ser Trp Trp Lys Asp Lys
                                    200
     67 His Asn Thr His His Ala Ala Pro Asn Val His Gly Glu Asp Pro Asp
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RAW SEQUENCE LISTING DATE: 06/28/2006
PATENT APPLICATION: US/10/583,084 TIME: 10:45:12

Input Set : A:\PTO.AMC.txt

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68
                           215
                                                220
70 Ile Asp Thr His Pro Leu Leu Thr Trp Ser Glu His Ala Leu Glu Met
                       230
                                           235
73 Phe Ser Asp Val Pro Asp Glu Glu Leu Thr Arg Met Trp Ser Arg Phe
                                       250
                   245
76 Met Val Leu Asn Gln Thr Trp Phe Tyr Phe Pro Ile Leu Ser Phe Ala
                                   265
79 Arg Leu Ser Trp Cys Leu Gln Ser Ile Met Phe Val Leu Pro Asn Gly
           275
                               280
82 Gln Ala His Lys Pro Ser Gly Ala Arq Val Pro Ile Ser Leu Val Glu
                           295
85 Gln Leu Ser Leu Ala Met His Trp Thr Trp Tyr Leu Ala Thr Met Phe
                       310
                                           315
88 Leu Phe Ile Lys Asp Pro Val Asn Met Ile Val Tyr Phe Leu Val Ser
                                       330
                   325
91 Gln Ala Val Cys Gly Asn Leu Leu Ala Ile Val Phe Ser Leu Asn His
92
               340
                                   345
94 Asn Gly Met Pro Val Ile Ser Lys Glu Glu Ala Val Asp Met Asp Phe
                               360
                                                    365
97 Phe Thr Lys Gln Ile Ile Thr Gly Arg Asp Val His Pro Gly Leu Phe
                           375
                                                380
100 Ala Asn Trp Phe Thr Gly Gly Leu Asn Tyr Gln Ile Glu His His Leu
101 385
                        390
                                            395
103 Phe Pro Ser Met Pro Arg His Asn Phe Ser Lys Ile Gln Pro Ala Val
                    405
                                         410
106 Glu Thr Leu Cys Lys Lys Tyr Gly Val Arg Tyr His Thr Thr Gly Met
                420
                                    425
109 Ile Glu Gly Thr Ala Glu Val Phe Ser Arg Leu Asn Glu Val Ser Lys
110
                                440
112 Ala Ala Ser Lys Met Gly Lys Ala Gln
113
        450
116 <210> SEQ ID NO: 2
117 <211> LENGTH: 1371
118 <212> TYPE: DNA
119 <213> ORGANISM: Mortierella alpina
121 <400> SEQUENCE: 2
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123 ctgaatgagg gcaagaagga tgccgaggca ccctttctga tgatcattga caacaaggtg 120
124 tacgatgtcc gcgagtttgt ccctgatcat cccggtggaa gtgtgattct cacgcacgtt 180
125 ggcaaggacg gcactgacgt ctttgacact ttccaccccg aggctgcttg ggagactctt 240
126 gccaactttt acgttggtga tattgatgag agcgatcgtg ccatcaagaa tgatgacttt 300
127 geggeegagg ttegeaaget gegeaeettg ttecagteee ttggetaeta egaetegtee 360
128 aaggcatact atgccttcaa ggtctcgttc aacctctgca tctggggctt gtcgactttc 420
129 attgttgcca agtggggcca gacctcgacc ctcgccaacg tgctctcggc tgcgctcttg 480
130 ggtctcttct ggcagcagtg cggatggttg gcgcacgact ttttgcacca ccaggtcttc 540
131 caggaccgtt tetggggtga tettttegge geettettgg gaggtgtetg ecagggttte 600
132 tegteeteet ggtggaagga caagcacaac acteaceaeg etgeteecaa egteeaegge 660
133 gaggateceg acattgacae teaceetetg ttgaeetgga gtgageatge tetggagatg 720
134 tteteggatg tteetgaega ggagetgaee egtatgtggt egegetteat ggteeteaae 780
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RAW SEQUENCE LISTING DATE: 06/28/2006 PATENT APPLICATION: US/10/583,084 TIME: 10:45:12

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06282006\J583084.raw

135 cagacotggt totacttocc cattotocg tittgcccgtc tgtcctggtg cotocagtcc 840 136 attatgtttg ttctgcccaa cggtcaggcc cacaagccct ctggagcgcg tgtgcccatt 900 137 tegttggteg ageagetgte tetggetatg caetggacet ggtacetege caecatgtte 960 138 ctgttcatta aggatcccgt caacatgatt gtgtactttt tggtgtcgca ggctgtttgc 1020 139 ggcaacttgt tggcgattgt gttctcgctc aaccacaacg gcatgcctgt gatctccaag 1080 140 gaggaagegg tegatatgga ettetteace aageagatea teaegggteg tgatgtteae 1140 141 cctggtctgt ttgccaactg gttcacgggt ggattgaact accagattga gcaccacttg 1200 142 ttcccttcga tgccccgcca caacttttca aagatccagc ctgctgtcga gactttgtgc 1260 143 aaaaagtacg gtgtccgata ccataccact ggtatgatcg agggaactgc agaggtcttt 1320 144 agccgtttga acgaggtctc caaggcggcc tccaaqatgg gcaaggcaca g 147 <210> SEQ ID NO: 3 148 <211> LENGTH: 318 149 <212> TYPE: PRT 150 <213> ORGANISM: Mortierella alpina 152 <400> SEQUENCE: 3 153 Met Glu Ser Ile Ala Gln Phe Leu Pro Ser Lys Met Pro Gln Asp Leu 10 156 Phe Ile Asp Leu Ala Arg Ala Ile Gly Val Gln Ala Ala Pro Tyr Val 20 25 159 Asp Pro Leu Glu Ala Ala Leu Val Ala Gln Ala Glu Lys Phe Pro 35 40 162 Thr Val Val His His Thr Arg Gly Phe Leu Val Ala Val Glu Ser Pro 163 50 55 165 Leu Ala Arg Glu Leu Pro Leu Met Asn Pro Phe His Val Leu Leu Ile 70 75 168 Ala Leu Ala Tyr Leu Val Thr Val Phe Val Gly Met Gln Ile Met Lys 85 171 Asn Phe Glu Arg Phe Glu Val Lys Thr Phe Ser Leu Phe His Asn Phe 172 100 105 174 Cys Leu Val Ser Ile Ser Ala Tyr Met Cys Gly Gly Ile Leu Tyr Glu 175 115 120 177 Ala Tyr Gln Ala Asn Tyr Gly Leu Phe Glu Asn Ala Ala Asp His Thr 130 135 180 Val Gln Gly Leu Pro Met Ala Lys Met Ile Trp Leu Phe Tyr Phe Ser 150 155 183 Lys Ile Met Glu Phe Val Asp Thr Met Ile Met Val Leu Lys Lys Asn 165 170 186 Asn Arg Gln Ile Ser Phe Leu His Val Tyr His His Ser Ser Ile Phe 180 185 190 189 Thr Ile Trp Trp Leu Val Thr Phe Val Ala Pro Asn Gly Glu Ala Tyr 200 192 Phe Ser Ala Ala Leu Asn Ser Phe Ile His Val Ile Met Tyr Gly Tyr 215 220 195 Tyr Phe Leu Ser Ala Leu Gly Phe Lys Gln Val Ser Phe Ile Lys Phe 235 198 Tyr Ile Thr Arg Ser Gln Met Thr Gln Phe Cys Met Met Ser Ile Gln 245 250 201 Ser Ser Trp Asp Met Tyr Ala Met Lys Val Leu Gly Arg Pro Gly Tyr 202 260 265

RAW SEQUENCE LISTING DATE: 06/28/2006 PATENT APPLICATION: US/10/583,084 TIME: 10:45:12

Input Set : A:\PTO.AMC.txt

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204 Pro Phe Phe Ile Thr Ala Leu Leu Trp Phe Tyr Met Trp Thr Met Leu
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                                280
                                                    285
207 Gly Leu Phe Tyr Asn Phe Tyr Arg Lys Asn Ala Lys Leu Ala Lys Gln
                            295
                                                300
210 Ala Lys Ile Asp Ala Ala Lys Glu Lys Ala Arg Lys Leu Gln
211 305
                        310
                                            315
214 <210> SEQ ID NO: 4
215 <211> LENGTH: 954
216 <212> TYPE: DNA
217 <213> ORGANISM: Mortierella alpina
219 <400> SEQUENCE: 4
220 atggagtega ttgegeaatt ceteceetea aagatgeege aagatetgtt tattgaeett 60
221 gcaagggcca teggtgtcca ggccgcaccc tatgtcgacc ctctcgaggc agcgcttgtg 120
222 gcccaggccg agaagttett ccccacggte gtteateaea egegeggett tttggtegeg 180
223 gtcgagtcac ccttggcccg tgagctgccc ttgatgaacc ccttccacgt gctgttgatc 240
224 gcgctcgctt acttggtcac ggtctttgtg ggcatgcaga tcatgaagaa ctttgaacgg 300
225 ttcgaggtca agacgttctc gctcttccac aacttttgtc tggtctcgat cagtgcctac 360
226 atgtgcggcg ggatcttgta cgaggcttac caggccaact atggactgtt tgagaacgcg 420
227 geogateata cegtecaggg tettectatg geoaagatga tetggetett etaettetee 480
228 aagatcatgg agtttgtcga caccatgatc atggtcctta agaagaacaa ccgccagatc 540
229 tegttettge aegtetacea ceacagetee atetteacea tetggtggtt ggteacettt 600
230 gttgcaccca atggtgaagc ctacttctcg gctgcgttga actcgttcat ccacgtgatc 660
231 atgtacggct actacttcct gtccgccttg ggcttcaagc aggtgtcgtt catcaagttc 720
232 tacatcacgc gttcgcagat gacgcagttc tgcatgatgt cgatccagtc ctcctgggac 780
233 atgtatgcca tgaaggtgct tggccgcccc ggatacccct tcttcatcac cgccctgctt 840
234 tggttctaca tgtggaccat gctcggactc ttctacaact tctacagaaa gaacgccaag 900
235 ttggccaagc aggccaagat cgatgctgcc aaggagaagg caaggaagtt gcag
238 <210> SEQ ID NO: 5
239 <211> LENGTH: 446
240 <212> TYPE: PRT
241 <213> ORGANISM: Mortierella alpina
243 <400> SEOUENCE: 5
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247 His Asn Thr Glu Asp Ser Leu Leu Leu Ala Ile Arg Gly Asn Val Tyr
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                                     25
250 Asp Val Thr Lys Phe Leu Ser Arg His Pro Gly Gly Thr Asp Thr Leu
253 Leu Leu Gly Ala Gly Arg Asp Val Thr Pro Val Phe Glu Met Tyr His
                             55
256 Glu Phe Gly Ala Ala Glu Ala Ile Met Lys Lys Tyr Tyr Val Gly Thr
                                             75
259 Leu Val Ser Asn Glu Leu Pro Ile Phe Pro Glu Pro Thr Val Phe His
260
262 Lys Thr Ile Lys Gly Arg Val Glu Ala Tyr Phe Lys Asp Arg Asn Met
263
                100
                                    105
265 Asp Ser Lys Asn Arg Pro Glu Ile Trp Gly Arg Tyr Ala Leu Ile Phe
            115
                                120
268 Gly Ser Leu Ile Ala Ser Tyr Tyr Ala Gln Leu Phe Val Pro Phe Val
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RAW SEQUENCE LISTING DATE: 06/28/2006 PATENT APPLICATION: US/10/583,084 TIME: 10:45:12

Input Set : A:\PTO.AMC.txt

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269
        130
                            135
271 Val Glu Arg Thr Trp Leu Gln Val Val Phe Ala Ile Ile Met Gly Phe
                        150
                                             155
274 Ala Cys Ala Gln Val Gly Leu Asn Pro Leu His Asp Ala Ser His Phe
                                         170
277 Ser Val Thr His Asn Pro Thr Val Trp Lys Ile Leu Gly Ala Thr His
                180
                                    185
280 Asp Phe Phe Asn Gly Ala Ser Tyr Leu Val Trp Met Tyr Gln His Met
            195
                                200
                                                     205
283 Leu Gly His His Pro Tyr Thr Asn Ile Ala Gly Ala Asp Pro Asp Val
                            215
                                                 220
286 Ser Thr Ser Glu Pro Asp Val Arg Arg Ile Lys Pro Asn Gln Lys Trp
                        230
                                             235
289 Phe Val Asn His Ile Asn Gln His Met Phe Val Pro Phe Leu Tyr Gly
                    245
                                         250
292 Leu Leu Ala Phe Lys Val Arg Ile Gln Asp Ile Asn Ile Leu Tyr Phe
                260
                                    265
                                                         270
295 Val Lys Thr Asn Asp Ala Ile Arg Val Asn Pro Ile Ser Thr Trp His
            275
                                280
298 Thr Val Met Phe Trp Gly Gly Lys Ala Phe Phe Val Trp Tyr Arg Leu
                            295
                                                 300
301 Ile Val Pro Met Gln Tyr Leu Pro Leu Ser Lys Val Leu Leu Leu Phe
                        310
                                            315
304 Thr Val Ala Asp Met Val Ser Ser Tyr Trp Leu Ala Leu Thr Phe Gln
                    325
                                         330
307 Ala Asn His Val Val Glu Glu Val Gln Trp Pro Leu Pro Asp Glu Asn
                                    345
310 Gly Ile Ile Gln Lys Asp Trp Ala Ala Met Gln Val Glu Thr Thr Gln
            355
                                360
313 Asp Tyr Ala His Asp Ser His Leu Trp Thr Ser Ile Thr Gly Ser Leu
        370
                            375
316 Asn Tyr Gln Ala Val His His Leu Phe Pro Asn Val Ser Gln His His
317 385
                        390
                                             395
319 Tyr Pro Asp Ile Leu Ala Ile Ile Lys Asp Thr Cys Ser Glu Tyr Lys
                    405
                                        410
322 Val Pro Tyr Leu Val Lys Asp Thr Phe Trp Gln Ala Phe Ala Ser His
                420
                                    425
325 Leu Glu His Leu Arg Val Leu Gly Leu Arg Pro Lys Glu Glu
           435
                                440
329 <210> SEQ ID NO: 6
330 <211> LENGTH: 1338
331 <212> TYPE: DNA
332 <213> ORGANISM: Mortierella alpina
334 <400> SEQUENCE: 6
335 atgggtacgg accaaggaaa aaccttcacc tggcaagaac tcgcggcgca taacaccgag 60
336 gacageetee ttttggetat eegtggeaat gtataegatg teacaaagtt ettgageegt 120
337 catcctggtg gaacggatac tctcttgctc ggagctggcc gagatgtcac tccggttttt 180
338 gagatgtacc acgagtttgg agctgcagag gctatcatga agaagtacta tgttggcaca 240
339 ctggtctcaa atgagttgcc catcttccca gagccaacgg tgttccataa gaccatcaag 300
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VERIFICATION SUMMARY

DATE: 06/28/2006 TIME: 10:45:13

PATENT APPLICATION: US/10/583,084

,004 IIME: IC

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\06282006\J583084.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

# Raw Sequence Listing before editing (for reference only)



**IFWP** 

RAW SEQUENCE LISTING DATE: 06/26/2006

PATENT APPLICATION: US/10/583,084 TIME: 13:40:03

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\06262006\J583084.raw

3 <110> APPLICANT: SUNTORY LIMITED

5 <120> TITLE OF INVENTION: Arachidonic acid-containing plants and use of the plants

7 <130> FILE REFERENCE: SU0423

C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/583,084

C--> 10 <141> CURRENT FILING DATE: 2006-06-15

12 <150> PRIOR APPLICATION NUMBER: JP 2003-419124

13 <151> PRIOR FILING DATE: 2003-12-17

15 <150> PRIOR APPLICATION NUMBER: JP 2004-097089

16 <151> PRIOR FILING DATE: 2004-03-29

18 <160> NUMBER OF SEQ ID NOS: 28

20 <170> SOFTWARE: PatentIn Ver. 2.1

# Does Not Comply Corrected Diskette Needed

#### ERRORED SEQUENCES

616 <210> SEQ ID NO: 28

617 <211> LENGTH: 20

618 <212> TYPE: DNA

619 <213> ORGANISM: Artificial Sequence

621 <220> FEATURE:

622 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer GLEr

624 <400> SEQUENCE: 28

625 cgacatcatg cagaactgtg

20

E--> 632/5E--> 635/5

E--> 637

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/583,084

DATE: 06/26/2006 TIME: 13:40:04

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\06262006\J583084.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:632 M:254 E: No. of Bases conflict, this line has no nucleotides. L:635 M:254 E: No. of Bases conflict, this line has no nucleotides. L:637 M:254 E: No. of Bases conflict, this line has no nucleotides.